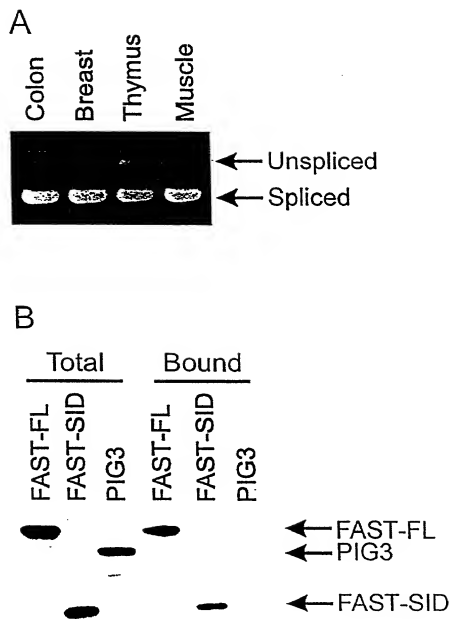


FIG. 1

ST-1	mgpcggsrlgpcgaespsqpp-----	21
ST-1	mrpcgslysgfcagsqyesveppslallssidqeqlpvatgqsynhsvqpw	51
ST-1	-----	21
ST-1	pqwpplsllyreggtwspdrsgmyglspgthegscctntheqpkdsmagdht	102
ST-1	---RRRRLRLRHDPEPYTYLAMEHLCGAAESRRRLHQAQIIRQQAQVFPF	69
ST-1	rsrRSKSGNRYRYYPFYSYLAMEHLCGNSPEKRLHQAQIIRQQAQVFPF	153
ST-1	FRDYEGWVDSERHNEGNNRCFRVPRDPAIPQAGNFWADYGLHFAEAL	120
ST-1	ENGDMVMDGDSERHNEGSSOCFKGLADFGAPQAGNFWADYGLHFAEAL	204
ST-1	RKONRHCGRWNGCARGAFAKDEGPYRHHGRPVrpsspppppsegfiks	171
ST-1	KRONRHCGRWNGCARGAFAKDEGPYRHHGRPVrpsspppppsegfiks	250
ST-1	lmgsggegawpplapqssvpagtgnsgeeaqptclpsserplwplcpl	222
ST-1	slslaedsqqtntggklntsfmidslhldlqadldasrnlengrjspav	301
ST-1	pgptrvegetvqggaigstlspcrawplllqgtavpggrssgghrasl	273
ST-1	amnmwssaplllythsketrnarsglstistystysssssistispvgfqk	352
ST-1	wgq-----	276
ST-1	eqeksgqrtqrvghpikrsredddcsttssdpdtnyspieppkkmplls	403
ST-1	FPSSVLPITYTNYMElappstscpcqcpstspaywgvapetrgppglc-	325
ST-1	cdpssvTKSVAENNAEpsvlgffhfrfttyynygpspytmppywgfphpt	454
ST-1	-----DLDALFQGVPPNCGSLQNVVYHPRDFAAGpggwllsw	362
ST-1	nsggdsprgpgqspldLDNMLRAMPNNKSLQDLTSHHGDEVHESflsqclg	505
ST-1	cg1-----	365
ST-1	sgsgpypsrqglm	518

FIG. 2



A

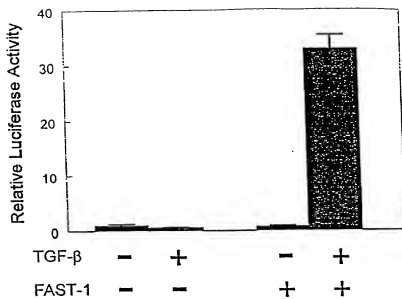
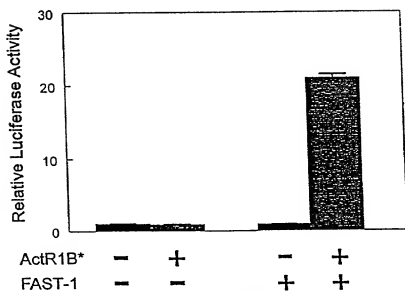


FIG. 3

B



C

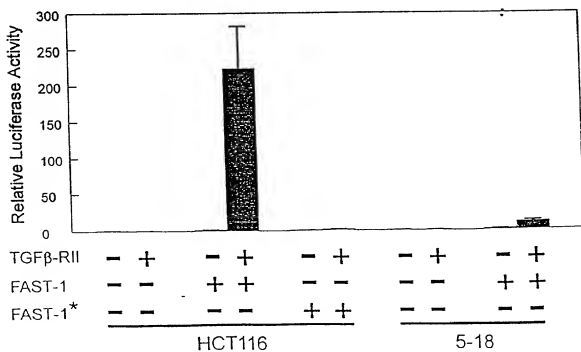
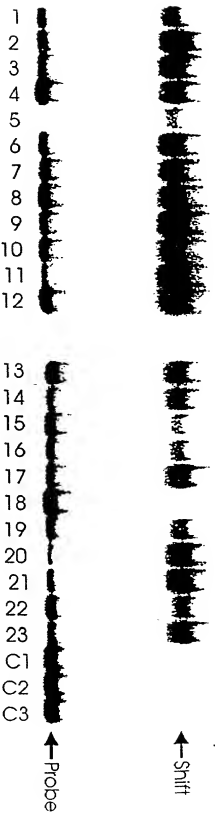


FIG. 4



B

		CONSENSUS									
		T	G	T	GT	GT	A	T	T		
G	41	41	11	0	100	0	54	47	0	0	70
A	23	36	17	0	0	0	5	0	100	0	64
T	23	23	64	100	0	100	52	0	0	100	11
C	11	0	5	0	0	0	0	0	0	0	5

